

EpiBuffer Examples

```
#####  
# DEPENDENCIES  
#####  
library(EpiBuffer) # SingleBuffer and SpatialBuffers functions  
library(dplyr)     # various helper functions  
library(tidyr)     # various helper functions  
library(fields)    # various helper functions  
library(ggplot2)   # plotting  
library(HDIInterval) # hdi function
```

Function to simulate outcomes, radii, exposure effect

```
sim_data_function <- function(exposure_indicator,  
                              unique_dists,  
                              unique_ps_dists,  
                              max_radius,  
                              x,  
                              w,  
                              q,  
                              v,  
                              beta_true,  
                              gamma_true,  
                              eta_true,  
                              tau_phi_true,  
                              rho_phi_true,  
                              sigma_epsilon_true,  
                              sim_setting,  
                              seed){  
  
  #####  
  # Global Settings  
  #####  
  n_ind <- nrow(x)  
  n_unique <- max(v)  
  
  spatial_cov <- tau_phi_true*tau_phi_true*exp(-rho_phi_true*unique_dists)  
  
  phi_true <- mnormt::rmnorm(n = 1,  
                             mean = rep(0.00, times = n_unique),  
                             varcov = spatial_cov)  
  
  # extract w, q, and phi for each individual based on location  
  v_w <- w[v, , drop = FALSE] # (n_ind x p_w)  
  v_q <- q[v, , drop = FALSE] # (n_ind x p_q)  
  v_phi <- phi_true[v, , drop = FALSE] # (n_ind x 1)
```

```

if(sim_setting == 0){ # no effect (theta = 0)

  radius_trans_true <- rep(0.00, times = n_ind)
  radius_true <- as.vector(max_radius*pnorm(radius_trans_true))
  theta_true <- rep(0.00, times = n_ind)

}

# True theta and radius (depends on sim setting)
if(sim_setting == 1){ # single radius, single theta

  radius_trans_true <- rep(gamma_true[1], times = n_ind)
  radius_true <- as.vector(max_radius*pnorm(radius_trans_true))
  theta_true <- rep(eta_true[1], times = n_ind)

}

if(sim_setting == 2){ # varying radii, single theta

  radius_trans_true <- (v_w%%gamma_true) + v_phi
  radius_true <- as.vector(max_radius*pnorm(radius_trans_true))
  theta_true <- rep(eta_true[1], times = n_ind) # theta = eta0 (pd = 0 setting)

}

if(sim_setting == 3){ # varying radii, varying theta

  radius_trans_true <- (v_w%%gamma_true) + v_phi
  radius_true <- max_radius*pnorm(radius_trans_true)

  # theta is coefficient on the radius (not on radius_trans!)
  theta_true <- v_q%%eta_true

}

# True exposure
exposure_true <- rep(NA, times = n_ind)
if(exposure_indicator == 0){ # counts
  v_exposure_dists <- unique_ps_dists[v, , drop = FALSE]
  exposure_true <- rowSums(v_exposure_dists < radius_true)
}
if(exposure_indicator == 2){ # presence/absence
  # exposed if at least one facility is within distance radius_true
  for(j in 1:n_ind){
    exposure_true[j] <- max(as.numeric(unique_ps_dists[v[j], ] <= radius_true[j]))
  }
}

#####
# Outcome
#####
# Simulate outcome from Gaussian distribution
# mu = x*\beta + \theta*exposure

```

```

mu <- x%%beta_true + theta_true*exposure_true
y <- rnorm(n = n_ind,
          mean = mu,
          sd = sigma_epsilon_true)

#####
# Return sim dat
#####
return(list(y = y,
            radius_true = radius_true,
            theta_true = theta_true,
            exposure_true = exposure_true))
}

```

Create inputs for sim data function

Specify simulation parameters

```

# Set seed for reproducibility
seed <- 3478

# Set all global simulation parameters
n_ind <- 500           # total number of outcome units
n_ind_unique <- 108    # total number of unique locations
m <- 100              # total number of exposure sources
max_radius <- 20       # maximum radius to consider
exposure_indicator <- 0 # 0: counts

# True parameter values
beta_true <- c(-0.95, 0.33)
tau_phi_true <- 1.00
rho_phi_true <- 3.07
gamma_true <- c(0.11, 0.07)
eta_true_single <- 0.90
sigma_epsilon_true <- 1.00

```

Simulate location and covariate information

```

set.seed(seed)

# Unique locations of outcome units
unique_locs <- data.frame(
  location_id = 1:n_ind_unique,
  x = runif(n_ind_unique, min = 0, max = max_radius*5),
  y = runif(n_ind_unique, min = 0, max = max_radius*5)
)

# Randomly assign outcome units to unique locations
ind_locs <- data.frame(
  individual_id = 1:n_ind,
  location_id = c(
    sample(1:n_ind_unique, size = n_ind_unique, replace = FALSE), # guarantee one per location

```

```

    sample(1:n_ind_unique, size = n_ind - n_ind_unique, replace = TRUE) # sample remaining
  )
)

# Sample locations of exposure sources (`ps`)
ps_locs <- data.frame(
  ps_id = 1:m,
  x = runif(m, min = 0, max = max_radius*5),
  y = runif(m, min = 0, max = max_radius*5)
)

#.....Create Location Matrices.....#

# v: index vector indicating unique location row in exposure_dists for each outcome unit
# length: n_ind
# NOTE: v maps each individual to a row in exposure_dists and w;
# rows must be ordered consistently with location indexing:
# row i corresponds to location_id[i], where location_id[v[j]] is the location for outcome unit j.
v <- match(ind_locs$location_id, unique_locs$location_id)

# Grid: knot locations for predictive process approximation
# here - same as unique locations (can be different or a subsample, as desired)
grid_coords <- unique_locs
grid_coords$location_id <- paste0("grid_", grid_coords$location_id)

all_locs <- rbind(unique_locs, grid_coords)

# unique_ps_dists: pairwise distance matrix for all unique locations and exposure sources
# only retain necessary exposure sources based on max_radius
unique_ps_dists <- fields::rdist(unique_locs[,c("x", "y")], ps_locs[,c("x", "y")])
unique_ps_dists <- unique_ps_dists[, !apply(unique_ps_dists > max_radius, 2, all)]

# unique_dists: pair-wise distance matrix for all unique locations
# dim: (n_ind_unique x n_ind_unique)
unique_dists <- fields::rdist(unique_locs[,c("x", "y")], unique_locs[,c("x", "y")])
unique_dists <- unique_dists / max(unique_dists)

# full_dists: pair-wise distance matrix for all unique locations and grid points
# dim: (n_ind_unique + n_grid) x (n_ind_unique + n_grid)
full_dists <- fields::rdist(all_locs[,c("x", "y")], all_locs[,c("x", "y")])
full_dists <- full_dists / max(full_dists) # scale to between 0 and 1

#.....Create Covariate Data.....#

# x: covariate matrix
# dim: nInd x p_x
x <- data.frame(
  x0 = rep(1, n_ind),
  x1 = rbinom(n_ind, size = 1, prob = 0.5) # a factor covariate
) %>% as.matrix()

# w: spatial covariate matrix (for SpatialBuffer)
# dim: n_unique x p_w

```

```

w <- data.frame(
  intercept = rep(1, n_ind_unique),
  w1 = rnorm(n_ind_unique, mean = 0, sd = 1)
) %>% as.matrix()

# q: spatial covariate matrix (for SpatialBuffer)
# dim: n_unique x p_q
q <- data.frame(
  intercept = rep(1, n_ind_unique),
  q1 = rnorm(n_ind_unique, mean = 0, sd = 1)
) %>% as.matrix()

```

Create simulated data sets from different settings

Recall -

- `sim_setting = 0`: no effect;
- `sim_setting = 1`: single radius, single effect;
- `sim_setting = 2`: varying radius, single effect;
- `sim_setting = 3`: varying radius, varying effect

```

# Simulated data where true dgp is single radius, single effect
sim_dat1 <- sim_data_function(exposure_indicator = exposure_indicator,
                             unique_dists = unique_dists,
                             unique_ps_dists = unique_ps_dists,
                             max_radius = max_radius,
                             x = x,
                             w = w,
                             q = q,
                             v = v,
                             beta_true = beta_true,
                             gamma_true = gamma_true,
                             eta_true = eta_true_single,
                             tau_phi_true = tau_phi_true,
                             rho_phi_true = rho_phi_true,
                             sigma_epsilon_true = sigma_epsilon_true,
                             sim_setting = 1,
                             seed = seed)

# Simulated data where true dgp is varying radius, single effect
sim_dat2 <- sim_data_function(exposure_indicator = exposure_indicator,
                             unique_dists = unique_dists,
                             unique_ps_dists = unique_ps_dists,
                             max_radius = max_radius,
                             x = x,
                             w = w,
                             q = q,
                             v = v,
                             beta_true = beta_true,
                             gamma_true = gamma_true,
                             eta_true = eta_true_single,
                             tau_phi_true = tau_phi_true,
                             rho_phi_true = rho_phi_true,

```

```
sigma_epsilon_true = sigma_epsilon_true,
sim_setting = 2,
seed = seed)
```

Fit Models

Set Global MCMC parameters

```
mcmc_samples <- 50000
burnin <- 30000
thin <- 2
keep_set <- seq(burnin, mcmc_samples, by = thin)
```

True DGP: Single Radius, Single Effect (sim_setting = 1)

FixedBuffer

```
print("Fitting SINGLEBUFFER")
```

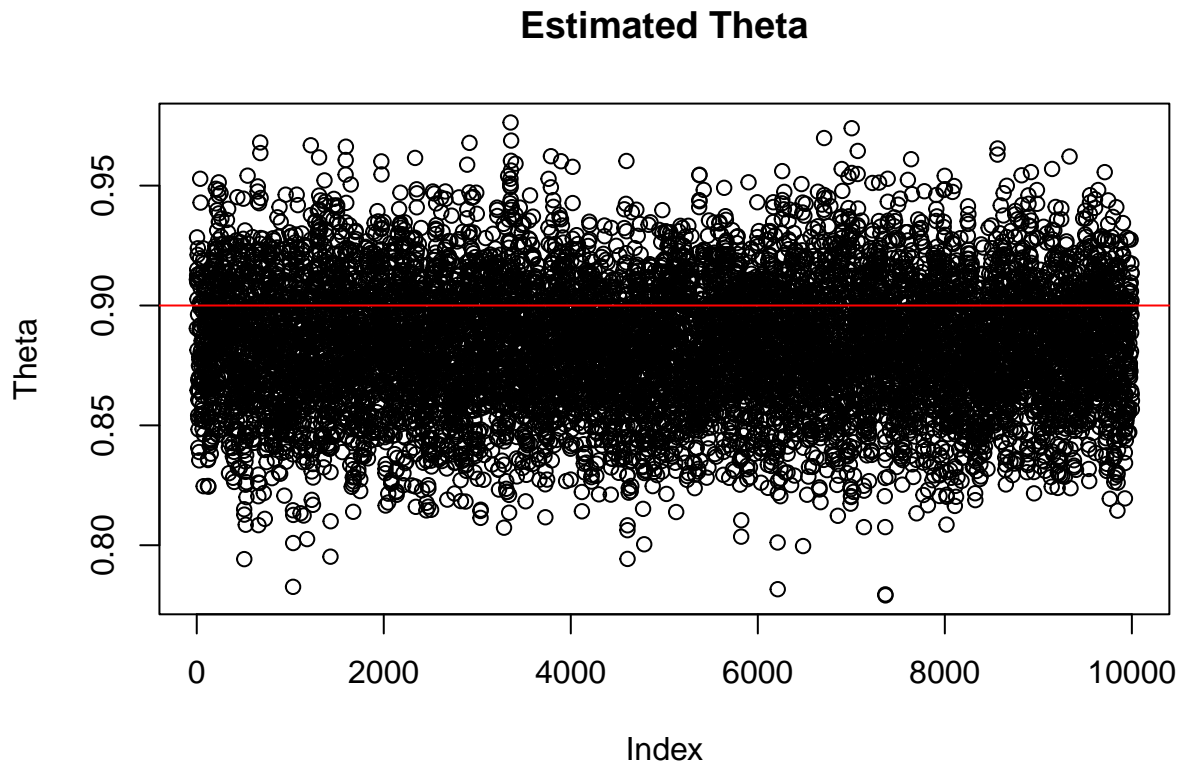
```
## [1] "Fitting SINGLEBUFFER"
```

```
metrop_var_radius <- 0.001
```

```
set.seed(seed)
res_fixed <- EpiBuffer::FixedBuffer(mcmc_samples = mcmc_samples,
                                   y = sim_dat1$y,
                                   x = x,
                                   q = matrix(q[,1], ncol = 1),
                                   v = v,
                                   radius = unique(sim_dat1$radius_true),
                                   exposure_definition_indicator = exposure_indicator,
                                   exposure_dists = unique_ps_dists,
                                   likelihood_indicator = 1,
                                   waic_info_indicator = 1,
                                   fitted_info_indicator = 1)
```

```
## Progress: 10%
## *****
## Progress: 20%
## *****
## Progress: 30%
## *****
## Progress: 40%
## *****
## Progress: 50%
## *****
## Progress: 60%
## *****
## Progress: 70%
## *****
## Progress: 80%
## *****
## Progress: 90%
## *****
```

```
## Progress: 100%
## *****
# Plot for theta
plot(res_fixed$theta[1, keep_set] / res_fixed$exposure_scale,
     main = "Estimated Theta",
     ylab = "Theta",
     xlab = "Index")
abline(h = unique(sim_dat1$theta_true), col = "red", lty = 1)
```



SingleBuffer

```
print("Fitting SINGLEBUFFER")

## [1] "Fitting SINGLEBUFFER"
metrop_var_radius <- 0.001

set.seed(seed)
res_single <- EpiBuffer::SingleBuffer(mcmc_samples = mcmc_samples,
                                     y = sim_dat1$y,
                                     x = x,
                                     q = matrix(q[,1], ncol = 1),
                                     v = v,
                                     radius_range = c(0.00, max_radius),
                                     exposure_definition_indicator = exposure_indicator,
                                     exposure_dists = unique_ps_dists,
```

```

metrop_var_radius = metrop_var_radius,
likelihood_indicator = 1,
waic_info_indicator = 1,
fitted_info_indicator = 1)

```

```

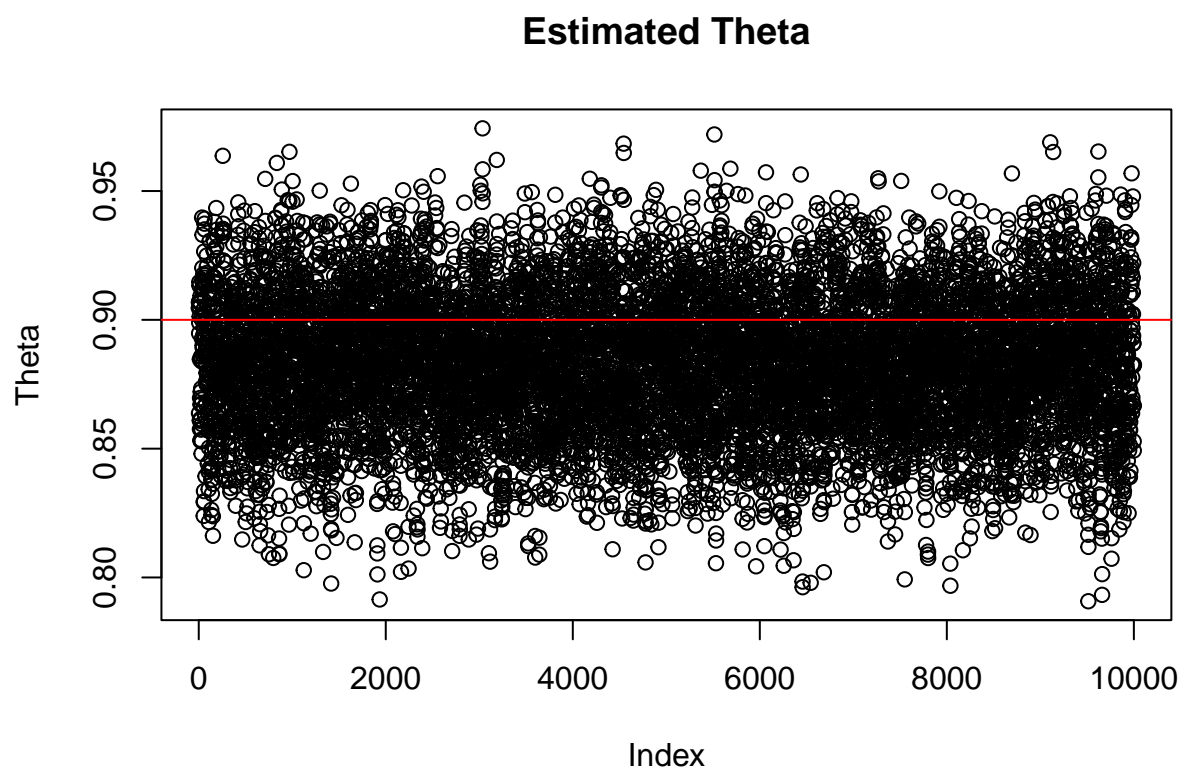
## Progress: 10%
## radius Acceptance: 12%
## *****
## Progress: 20%
## radius Acceptance: 12%
## *****
## Progress: 30%
## radius Acceptance: 12%
## *****
## Progress: 40%
## radius Acceptance: 12%
## *****
## Progress: 50%
## radius Acceptance: 12%
## *****
## Progress: 60%
## radius Acceptance: 12%
## *****
## Progress: 70%
## radius Acceptance: 12%
## *****
## Progress: 80%
## radius Acceptance: 12%
## *****
## Progress: 90%
## radius Acceptance: 12%
## *****
## Progress: 100%
## radius Acceptance: 12%
## *****

```

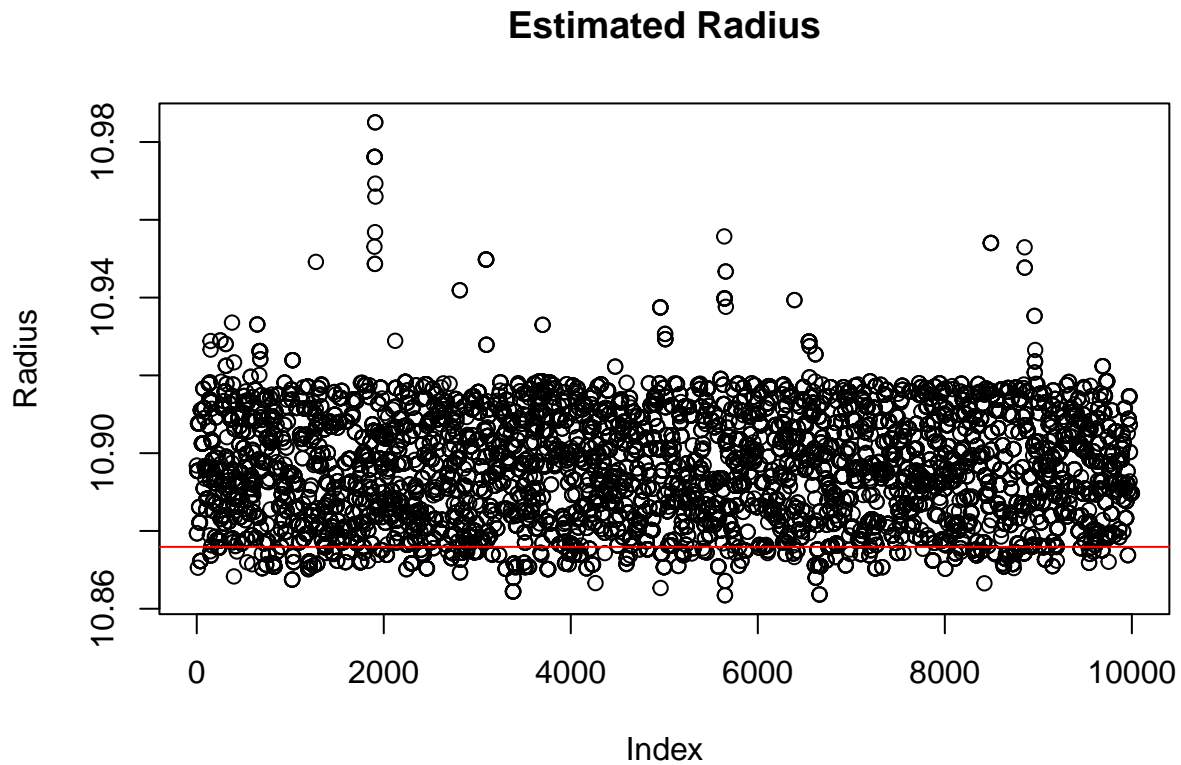
```

# Plot for theta
plot(res_single$theta[1, keep_set] / res_single$exposure_scale,
     main = "Estimated Theta",
     ylab = "Theta",
     xlab = "Index")
abline(h = unique(sim_dat1$theta_true), col = "red", lty = 1)

```

```
# Plot for radius
plot(res_single$radius[keep_set],
     main = "Estimated Radius",
     ylab = "Radius",
     xlab = "Index")
abline(h = unique(sim_dat1$radius_true), col = "red", lty = 1)
```



True DGP: Varying Radius, Varying Effect (sim_setting = 2)

SpatialBuffer

```
print("Fitting SPATIALBUFFER")
```

```
## [1] "Fitting SPATIALBUFFER"
```

```
metrop_var_gamma <- 0.01
```

```
metrop_var_phi_star <- 0.2
```

```
metrop_var_tau_phi <- 0.85
```

```
metrop_var_rho_phi <- 1.0
```

```
set.seed(seed)
```

```
res_spatial1 <- EpiBuffer::SpatialBuffers(mcmc_samples = mcmc_samples,
  y = sim_dat2$y,
  x = x,
  w = w,
  q = matrix(q[,1], ncol = 1),
  v = v,
  radius_range = c(0.00, max_radius),
  exposure_definition_indicator = exposure_indicator,
  exposure_dists = unique_ps_dists,
  full_dists = full_dists,
  metrop_var_gamma = rep(metrop_var_gamma,
    times = ncol(w)),
```

```

metrop_var_phi_star = rep(metrop_var_phi_star,
                           times = length(v)),
metrop_var_tau_phi = metrop_var_tau_phi,
metrop_var_rho_phi = metrop_var_rho_phi,
likelihood_indicator = 1,
waic_info_indicator = 1,
fitted_info_indicator = 1)

```

```

## Progress: 10%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 65%
## rho_phi Acceptance: 17%
## *****
## Progress: 20%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 65%
## rho_phi Acceptance: 17%
## *****
## Progress: 30%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 64%
## rho_phi Acceptance: 17%
## *****
## Progress: 40%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 43%
## tau_phi Acceptance: 65%
## rho_phi Acceptance: 17%
## *****
## Progress: 50%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 66%
## rho_phi Acceptance: 17%
## *****
## Progress: 60%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 66%

```

```

## rho_phi Acceptance: 17%
## *****
## Progress: 70%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 66%
## rho_phi Acceptance: 17%
## *****
## Progress: 80%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 66%
## rho_phi Acceptance: 17%
## *****
## Progress: 90%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 65%
## rho_phi Acceptance: 17%
## *****
## Progress: 100%
## gamma Acceptance (min): 4%
## gamma Acceptance (max): 5%
## phi_star Acceptance (min): 6%
## phi_star Acceptance (max): 44%
## tau_phi Acceptance: 65%
## rho_phi Acceptance: 17%
## *****

# Extract output and plot results
radius_df <- data.frame(t(res_spatial1$radius)) # Rows = MCMC iterations; columns = outcome units
radii_thinned <- radius_df[keep_set,]
radius_post_summary <- radii_thinned %>%
  dplyr::summarise(across(everything(), list( # apply to each column (individual)
    posterior.median = ~median(.),
    posterior.mean = ~mean(.),
    hdi.lower95 = ~hdi(., credMass = 0.95)[["lower"]],
    hdi.upper95 = ~hdi(., credMass = 0.95)[["upper"]],
    ci.lower95 = ~quantile(., 0.025),
    ci.upper95 = ~quantile(., 0.975)
  ))) %>%
  pivot_longer(cols = everything(), names_to = c("cluster", ".value"), names_sep = "_")
radius_post_summary$radius_true <- sim_dat2$radius_true

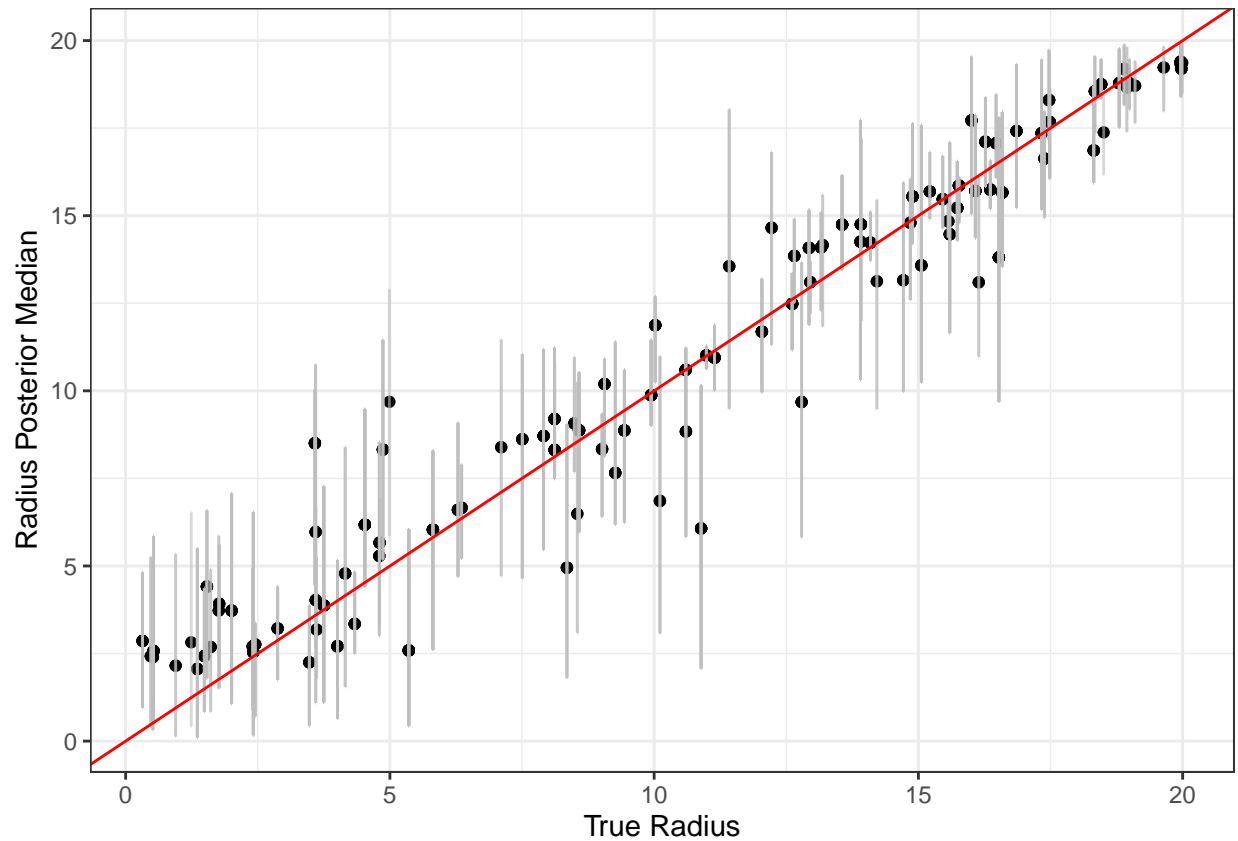
theta_df <- data.frame(t(res_spatial1$theta))
theta_df <- theta_df/res_spatial1$exposure_scale # unscale thetas
theta_thinned <- theta_df[keep_set, ]

```

```

ggplot(radius_post_summary,
       aes(x = radius_true, y = posterior.median)) +
  geom_point() +
  geom_errorbar(aes(ymin = hdi.lower95, ymax = hdi.upper95), # Add credible intervals
               width = 0.01,
               color = "grey", alpha = 0.6) +
  xlab("True Radius") +
  ylab("Radius Posterior Median") +
  theme_bw() +
  geom_abline(slope = 1, intercept = 0, color = "red")

```



```

ggplot(theta_thinned, aes(x = `X1`)) +
  geom_histogram(bins = 40) +
  geom_vline(xintercept = sim_dat2$theta_true, color = "red") +
  xlab("Theta Posterior Samples") +
  theme_bw() +
  theme(axis.text.x=element_text(size=12))

```

